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Set	Items	Description
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S1	84	RAGE AND REVIEW
S2	44	S1 AND RECEPTOR
S3	1	S2 AND SPLICE
S4	1	S1 AND SPLICE
S5	1	S1 AND ALTERNATIVE
S6	686	RECEPTOR(3W)ADVANCED() GLYCATION
S7	14	S6 AND SPLICE

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3/7/1

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18336797 BIOSIS NO.: 200510031297

Roles of the ~~receptor~~ for advanced glycation endproducts in
diabetes-induced vascular injury

AUTHOR: Yonekura Hideto (Reprint); Yamamoto Yasuhiko; Sakurai Shigeru;
Watanabe Takuo; Yamamoto Hiroshi

AUTHOR ADDRESS: Kanazawa Univ, Grad Sch Med Sci, Dept Biochem and Mol Vasc
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JOURNAL: Journal of Pharmacological Sciences 97 (3): p305-311 MAR 05 2005
ISSN: 1347-8613

DOCUMENT TYPE: Article; Literature Review

RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: Diabetic patients have shorter life span and poorer Quality of Life mainly due to diabetic vascular complications. Recent in vitro and in vivo studies have shown that advanced glycation endproducts (AGE) account for diabetic vascular complications through their engagement of the ~~receptor~~ for AGE (~~RAGE~~). In this ~~review~~, we summarize our recent studies on the roles of the AGE-~~RAGE~~ system in diabetes-induced vascular injury. In vitro experiments showed that AGE engagement of PAGE leads to changes in endothelial cells (EC) and pericytes, which are characteristic of diabetic microangiopathy. Diabetic ~~RAGE~~ transgenic mice that overexpress ~~RAGE~~ in vascular cells exhibited the exacerbation of the indices of nephropathy and retinopathy, and this was prevented by the inhibition of AGE formation. ~~RAGE~~ overexpression also caused calcium handling impairment in cardiac myocytes. In contrast to the ~~RAGE~~-overexpressing mice, diabetic ~~RAGE~~ knockout mice showed marked improvement of nephropathy. We found that human vascular cells express a novel ~~splice~~ variant coding for a soluble ~~RAGE~~ protein and named it endogenous secretory ~~RAGE~~ (esRAGE). The esRAGE neutralizes AGE actions on EC and is present in human sera. Individual variations in circulating esRAGE could be a determinant for individual differences in susceptibility or

resistance to the development of diabetic vascular complications. The AGE-~~AGE~~ system should be, therefore, a candidate molecular target for overcoming diabetic vascular complications.

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4/7/1

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18336797 BIOSIS NO.: 200510031297

Roles of the receptor for advanced glycation endproducts in diabetes-induced vascular injury

AUTHOR: Yonekura Hideto (Reprint); Yamamoto Yasuhiko; Sakurai Shigeru; Watanabe Takuo; Yamamoto Hiroshi

AUTHOR ADDRESS: Kanazawa Univ, Grad Sch Med Sci, Dept Biochem and Mol Vasc Biol, Kanazawa, Ishikawa 9208640, Japan**Japan

AUTHOR E-MAIL ADDRESS: hyone@med.kanazawa-u.ac.jp

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ABSTRACT: Diabetic patients have shorter life span and poorer Quality of Life mainly due to diabetic vascular complications. Recent in vitro and in vivo studies have shown that advanced glycation endproducts (AGE) account for diabetic vascular complications through their engagement of the receptor for AGE (~~AGE~~). In this ~~review~~, we summarize our recent studies on the roles of the AGE-~~AGE~~ system in diabetes-induced vascular injury. In vitro experiments showed that AGE engagement of ~~AGE~~ leads to changes in endothelial cells (EC) and pericytes, which are characteristic of diabetic microangiopathy. Diabetic ~~AGE~~ transgenic mice that overexpress ~~AGE~~ in vascular cells exhibited the exacerbation of the indices of nephropathy and retinopathy, and this was prevented by the inhibition of AGE formation. ~~AGE~~ overexpression also caused calcium handling impairment in cardiac myocytes. In contrast to the ~~AGE~~-overexpressing mice, diabetic ~~AGE~~ knockout mice showed marked improvement of nephropathy. We found that human vascular cells express a novel ~~splice~~ variant coding for a soluble ~~AGE~~ protein and named it endogenous secretory ~~AGE~~ (esRAGE). The esRAGE neutralizes AGE actions on EC and is present in human sera. Individual variations in circulating esRAGE could be a determinant for individual differences in susceptibility or resistance to the development of diabetic vascular complications. The AGE-~~AGE~~ system should be, therefore, a candidate molecular target for overcoming diabetic vascular complications.

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7/7/1

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0019461037 BIOSIS NO.: 200700120778

Lentivirus gene transfer of the endogenous circulating RAGE ~~splice~~
form blocks mechanisms leading to atherosclerosis
AUTHOR: Hudson Barry I (Reprint); Harja Evis; Arriero Maria; Nikolla Zana;
Schmidt Anna Marie
AUTHOR ADDRESS: Columbia Univ, New York, NY 10027 USA**USA
JOURNAL: Circulation 114 (18, Suppl. S): p25-26 OCT 31 2006 2006
CONFERENCE/MEETING: 79th Annual Scientific Session of the
American-Heart-Association Chicago, IL, USA November 12 -15, 2006;
20061112
SPONSOR: Amer Heart Assoc
ISSN: 0009-7322
DOCUMENT TYPE: Meeting; Meeting Abstract
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: The ~~Receptor~~ for ~~Advanced~~ ~~Glycation~~
End-products (RAGE) is a strong candidate gene for the development of
atherosclerosis. We have previously shown that pharmacological blockade
of RAGE (recombinant soluble RAGE) or genetic deletion of RAGE suppresses
the development of vascular disease in animal models. Further, in human
subjects the endogenous circulating levels of RAGE are lower in plasma of
subjects with coronary artery disease, suggesting an innate protective
role for circulating soluble (s)RAGE. Characterization of cardiovascular
tissue and cells by RT-PCR revealed the presence of RAGE and the
production of sRAGE by alternative splicing resulting from inclusion of
intron 9 (RAGEint9) in normal human heart and in endothelial (EC), aortic
smooth muscle (AoSMC) and monocyte/macrophage (MP) cells. Using an
antibody generated against the unique C-terminus protein ~~splice~~
region of RAGEint9, we detected RAGEint9 in lysates and cultured media
from EC, aortic (Ao) SMC and macrophages. Transfection of RAGEint9
suppressed RAGE-ligand S100B-stimulation of MMP-9 activity and levels by
similar to 2-3 and similar to 3-4-fold respectively, compared to
wild-type RAGE expressing cells (p < 0.0001). Upon S100B incubation,
RAGEint9- vs. RAGE expressing cells generated significantly decreased
levels of IL-6 in the supernatant (similar to 1.6-fold; p < 0.0001). We
generated lentiviral constructs to enable gene transfer of RAGEint9 into
primary human aortic SMCs and ECs. Lentivirus transduction of RAGEint9
into human AoSMCs, reduced S100B-stimulated levels of MCP-1 by similar to
2-fold compared to cells expressing full-length RAGE (p < 0.05). Further,
the migration of AoSMCs was blocked in RAGEint9 transduced cells in
response to RAGE ligand stimulation. Finally, in ECs transduced with RAGE
/ RAGEint9 lentivirus, permeability assays were performed. The gene
transfer of RAGEint9 into ECs blocked RAGE-ligand dependent endothelial
permeability by 50%, p < 0.001, the hallmark of atherosclerosis. We
conclude that RAGEint9 gene therapy or endogenous stimulated expression
may represent a novel therapeutic modality to enhance protection against
atherosclerotic disease.

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0019458768 BIOSIS NO.: 200700118509
Endogenous secretory ~~receptor~~ for ~~advanced~~ ~~glycation~~ end
products in non-small cell lung carcinoma
AUTHOR: Kobayashi Seichi; Kubo Hiroshi (Reprint); Suzuki Takashi; Ishizawa

Kota; Yamada Mitsuhiro; He Mei; Yamamoto Yasuhiko; Yamamoto Hiroshi;
Sasano Hironobu; Sasaki Hidetada; Suzuki Satoshi
AUTHOR ADDRESS: Tohoku Univ, Sch Med, Dept Geriatr and Resp Med, 1-1
Seiryomachi, Sendai, Miyagi 9808574, Japan**Japan
AUTHOR E-MAIL ADDRESS: hkubo@geriat.med.tohoku.ac.jp
JOURNAL: American Journal of Respiratory and Critical Care Medicine 175 (2
): p184-189 JAN 15 2007 2007
ISSN: 1073-449X
DOCUMENT TYPE: Article
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: Rationale: The **receptor** for **advanced** **glycation** end products is a multiligand receptor that plays an important role in regulating the invasiveness and metastatic potential of cancer cells. A recently discovered novel **splice** variant, the endogenous secretory **receptor** for **advanced** **glycation** end products, mediates the **receptor** for **advanced** **glycation** end-product-associated cell responses by functioning as a decoy receptor.Objectives: To evaluate the expression pattern of endogenous secretory **receptor** for **advanced** **glycation** end products in non-small cell lung carcinoma, and analyze its impact on prognosis.Methods: We performed immunohistochemical evaluation in 182 non-small cell lung carcinoma surgical specimens. The effect of an overexpressed receptor in cancer cell proliferation was also evaluated.Measurements and Main Results: The endogenous secretory **receptor** for **advanced** **glycation** end-product expression in cytoplasm was reduced or absent in 137 of the 182 (75%) carcinomas in contrast to normal lung tissues. mRNA expression was also suppressed in cancer cells. Overexpression of the secretory receptor in lung cancer cell lines had an inhibitory effect on cell proliferation, suggesting the reduced receptor expression accelerated tumor growth. Among patients with low expression of the cytoplasmic secretory receptor, the overall survival rate was significantly lower than that of patients with normal expression ($p = 0.0003$). This association was most prominent in TNM stage I patients ($p = 0.0001$). In a multivariate analysis, endogenous secretory receptor immunoreactivity was an independent prognostic factor with a relative risk of 3.1.Conclusions: The cytoplasmic endogenous secretory **receptor** for **advanced** **glycation** end-product expression has the potential to be a prognostic factor for predicting the outcome of curative surgery in patients with non-small cell lung carcinoma.

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19065460 BIOSIS NO.: 200600410855

Identification of mouse orthologue of endogenous secretory **receptor** for **advanced** **glycation** end-products: structure, function and expression

AUTHOR: Harashima Ai; Yamamoto Yasuhiko (Reprint); Cheng Chunmei; Tsuneyama Koichi; Myint Khin Mar; Takeuchi Akihiko; Yoshimura Kazunobu; Li Hui; Watanabe Takuo; Takasawa Shin; Okamoto Hiroshi; Yonekura Hideto; Yamamoto Hiroshi

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AUTHOR E-MAIL ADDRESS: yasuyama@med.kanazavva-u.ac.jp
JOURNAL: Biochemical Journal 396 (Part 1): p109-115 MAY 15 2006 2006
ISSN: 0264-6021
DOCUMENT TYPE: Article
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: The cell-surface RAGE [%%receptor%% for AGE (%%advanced%%
%%glycation%% end-products)] is associated with the development of
diabetic vascular complications, neurodegenerative disorders and
inflammation. Recently, we isolated a human RAGE %%splice%% variant,
which can work as a decoy receptor for RAGE ligands, and named it esRAGE
(endogenous secretory RAGE). In the present study, we have isolated the
murine equivalent of esRAGE from brain polysomal poly(A) (+)
(polyadenylated) RNA by RT (reverse transcription)-PCR cloning. The mRNA
was generated by alternative splicing, and it encoded a 334-amino-acid
protein with a signal sequence, but lacking the transmembrane domain. A
transfection experiment revealed that the mRNA was actually translated
as deduced to yield the secretory protein working as a decoy in
AGE-induced NF-kappa B (nuclear factor kappa B) activation. RT-PCR and
immunoblotting detected esRAGE mRNA and protein in the brain, lung,
kidney and small intestine of wild-type mice, but not of RAGE-null mice.
The esRAGE expression was increased in the kidney of diabetic wild-type
mice. The present study has thus provided an animal orthologue of esRAGE
for clarification of its roles in health and disease.

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18991676 BIOSIS NO.: 200600337071

Alternative splicing of the RAGE gene: Analysis and characterization in
humans and mice

AUTHOR: Hudson Barry Ian (Reprint); Arriero Maria; Yang Hojin; Moser
Bernhard; Harja Evis; Schmidt Ann Marie

AUTHOR ADDRESS: Columbia Univ, New York, NY 10032 USA**USA

JOURNAL: FASEB Journal 20 (5, Part 2): pA1081 MAR 7 2006 2006

CONFERENCE/MEETING: Experimental Biology 2006 Meeting San Francisco, CA,
USA April 01 -05, 2006; 20060401

SPONSOR: Amer Assoc Anatomists

Amer Physiol Soc

Amer Soc Biochem & Mol Biol

Amer Soc Investigat Pathol

Amer Soc Nutr

Amer Soc Pharmacol & Expt Therapeut

ISSN: 0892-6638

DOCUMENT TYPE: Meeting; Meeting Abstract

RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: The alternative splicing of pre-mRNAs is a critical mechanism in
genomic complexity, disease, and development. Studies on the
%%Receptor%% for %%Advanced%% %%Glycation%% End-products (RAGE)
indicate this gene undergoes a variety of %%splice%% events; however,
no studies have analyzed the tissue distribution or compared evolutionary
differences between species of RAGE isoforms. We recently reported on the

analysis of RAGE splicing in lung by a novel method we termed the Rapid Identification, Characterization and Expression of %%%splice%%% forms (RICE-%%%SPLICE%%%). We applied this method to cDNA from multiple human and mouse tissue extracts for RAGE including lung, heart, brain, kidney, pancreas and placenta and compared the %%%splice%%% forms found between these two species. In humans, numerous %%%splice%%% variants were identified ranging from deletion of multiple exons (entire or in part) and intronic inclusions. These variants were predicted to change the structure and function of RAGE since they resulted in alterations in the ligand-binding domain, extensive removal of the extracellular region, production of novel proteins, and the production of soluble forms lacking the transmembrane region. In contrast, in mice, fewer %%%splice%%% variants were identified, mainly resulting in forms lacking the transmembrane region. Comparison of %%%splice%%% variants between humans and mice revealed homologous regions in the RAGE gene which undergo splicing and similar %%%splice%%% forms. Further analysis of these forms revealed the biological significance of the conserved %%%splice%%% variants. In conclusion, we have identified differences in the tissue and cross-species distribution of RAGE %%%splice%%% forms which further expands the biological repertoire of this receptor in health and disease.

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18679332 BIOSIS NO.: 200600024727

%%%Receptor%%% for %%advanced%%% %%%glycation%%% end products is a promising target of diabetic nephropathy

BOOK TITLE: Annals of the New York Academy of Sciences

AUTHOR: Yamamoto Yasuhiko; Doi Toshio; Kato Ichiro; Shinohara Harumichi; Sakurai Shigeru; Yonekura Hideto; Watanabe Takuo; Myint Kihn Mar; Harashima Ai; Takeuchi Masayoshi; Takasawa Shin; Okamoto Hiroshi; Hashimoto Noriyoshi; Asano Masahide; Yamamoto Hiroshi (Reprint)

BOOK AUTHOR/EDITOR: Baynes JW (Editor); Monnier VM (Editor); Ames JM (Editor); Thorpe SR (Editor)

AUTHOR ADDRESS: Kanazawa Univ, Grad Sch Med Sci, Dept Biochem and Mol Vasc Biol, 13-1 Takara Machi, Kanazawa, Ishikawa 9208640, Japan**Japan

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SERIES TITLE: ANNALS OF THE NEW YORK ACADEMY OF SCIENCES 1043 p562-566 2005

BOOK PUBLISHER: NEW YORK ACAD SCIENCES, 2 EAST 63RD ST, NEW YORK, NY 10021 USA

CONFERENCE/MEETING: 8th International Symposium on the Maillard Reaction Charleston, SC, USA August 29 -September 01, 2004; 20040829

SPONSOR: Asahi Kasei Pharma

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Univ S Carolina, Res & Hlth Sci
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ISSN: 0077-8923_(print) ISBN: 1-57331-531-1 (H)

DOCUMENT TYPE: Book Chapter; Meeting

RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: Advanced glycation end products (AGEs) and the receptor for AGE (RAGE) interactions have been implicated in the development of diabetic vascular complications, which cause various disabilities and shortened life expectancy, and reduced quality of life in patients with diabetes. Diabetes-induced RAGE-overexpressing transgenic mice exhibited the exacerbation of the indices of nephropathy, and this was prevented by the inhibition of AGE formation. We also created RAGE-deficient mice by homologous recombination. They showed marked amelioration of diabetic nephropathy as compared with wild-type mice. Through an analysis of vascular polysomal poly(A)(+) RNA, we identified a novel **splice** variant coding for a soluble RAGE protein and named it endogenous secretory RAGE (esRAGE). esRAGE was able to protect AGE-induced vascular cell injuries as a decoy receptor and was actually detected in human circulation. We conclude that RAGE plays an active role in the development of diabetic vascular complications, especially nephropathy, and is a promising target for overcoming this disease. The esRAGE, an endogenous decoy receptor, may be related to individual variations in resistance to the development of diabetic vascular complications.

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18648479 BIOSIS NO.: 200510342979

Expression profiling of endogenous secretory **receptor** for **advanced** **glycation** end products in human organs

AUTHOR: Cheng Chunmei; Tsuneyama Koichi (Reprint); Kominami Rieko; Shinohara Harumichi; Sakurai Shigeru; Yonekura Hideto; Watanabe Takuo; Takano Yasuo; Yamamoto Hiroshi; Yamamoto Yasuhiko

AUTHOR ADDRESS: Toyama Med and Pharmaceut Univ, Dept Pathol, COE Program 21, 2630 Sugitani, Toyama 9300194, Japan**Japan

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JOURNAL: Modern Pathology 18 (10): p1385-1396 OCT 2005 2005

ISSN: 0893-3952

DOCUMENT TYPE: Article

RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: The **receptor** for **advanced** **glycation** end

products (RAGE) is a cell surface multiligand receptor of the immunoglobulin superfamily, which participates in physiological and pathological processes such as neuronal development, diabetes, inflammation, neurodegenerative disorders, and cancer. A novel ~~splice~~ variant of RAGE- endogenous secretory decoy form (esRAGE) was recently identified and is thought to be a prospective candidate to modify these RAGE- associated conditions. Here, we investigated the expression and distribution of esRAGE and RAGE proteins with domain-specific antibodies. We studied a wide variety of adult normal human preparations obtained from surgical and autopsy specimens using a tissue microarray technique. The results revealed that esRAGE was widely distributed and we classified its expression into four patterns. In pattern A, the cytoplasm is stained diffusely in neurons, vascular endothelium, pneumocytes, mesothelium, pancreatic beta cells, and macrophages/ monocytes. In pattern B, dot- like granules are stained in the supranuclear regions facing the luminal surface of the bile ducts, salivary glands, digestive tracts, renal tubules, prostate, skin, thyroid, and bronchioles. Pattern C is represented by diffuse staining in the stromal area of the arterial walls. Pattern D shows diffuse and strong staining of secreted materials such as thyroidal colloid, crystals in renal tubular lumen, and glandular lumen in prostate. This study provides, for the first time, a histopathological basis for understanding the physiological roles of esRAGE in humans, and will contribute to elucidating the participation of esRAGE in pathological processes and to exploring novel diagnostic and therapeutic concepts.

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18631205 BIOSIS NO.: 200510325705

A novel method for the detailed analysis of gene ~~splice~~ variants

AUTHOR: Hudson Barry Ian (Reprint); Carter Angela M; Harja Evis; Arriero

Maria; Yang Hojin; Moser Bernhard; Grant Peter J; Schmidt Ann Marie

AUTHOR ADDRESS: Columbia Univ, Dept Surg, New York, NY 10032 USA**USA

JOURNAL: FASEB Journal 19 (4, Suppl. S, Part 1): pA855 MAR 4 2005 2005

CONFERENCE/MEETING: Experimental Biology 2005 Meeting/35th International Congress of Physiological Sciences San Diego, CA, USA March 31.-April 06, 2005; 20050331

SPONSOR: Amer Assoc Anatomists

Amer Assoc Immunologists

Amer Physiol Soc

Amer Soc Biochem & Mol Biol

Amer Soc Investigat Pathol

Amer Soc Nutr Sci

Amer Soc Pharmacol & Expt Therapeut

Int Union Physiol Sci

ISSN: 0892-6638

DOCUMENT TYPE: Meeting; Meeting Abstract

RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: Alternative splicing of pre-mRNAs is a critical mechanism to produce a diverseproteome. Further development of methodology is needed for the thorough identification of ~~splice~~ variants produced in physiological and diseased states.The objective of this study was to

Print

perform detailed analysis of gene ~~splice~~ variants using a novel method. As an example, we have analyzed the gene encoding the ~~Receptor~~ for ~~Advanced~~ ~~Glycation~~ End-products (RAGE) for ~~splice~~ variants. ~~RAGE~~ variants were PCR-amplified from lung cDNA, cloned and analyzed by PCR-restriction digestion. Transfection of the characterized clones, into cells allowed the identification of the resulting protein. We have called this method the Rapid Identification, Characterization and Expression of ~~Splice~~ forms (RICE-~~SPLICE~~). Numerous ~~splice~~ variants of RAGE were identified ranging from deletion of multiple exons (entire or in part) and intronic inclusion. The variants were predicted to change the structure and function of RAGE since they resulted in alterations in the ligand-binding domain; extensive removal of the extracellular region; production of novel proteins and the production of soluble forms lacking the transmembrane region. Variants expressed in vitro were shown to produce functional proteins. Application of this method to other genes would facilitate detailed characterization of ~~splice~~ forms relevant to homeostasis and disease.

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18457565 BIOSIS NO.: 200510152065

Modulation of soluble ~~receptor~~ for ~~advanced~~ ~~glycation~~ end products by angiotensin-converting enzyme-1 inhibition in diabetic nephropathy

AUTHOR: Forbes Josephine M (Reprint); Thorpe Suzanne R; Thallas-Bonke Vicki ; Pete Josefa; Thomas Merlin C; Deemer Elizabeth R; Bassal Sahar; El-Osta Assam; Long David M; Panagiotopoulos Sianna; Jerums George; Osicka Tanya M; Cooper Mark E

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JOURNAL: Journal of the American Society of Nephrology 16 (8): p2363-2372

AUG 05 2005

ISSN: 1046-6673

DOCUMENT TYPE: Article

RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: Recent studies have identified that first-line renoprotective agents that interrupt the renin-angiotensin system not only reduce BP but also can attenuate advanced glycation end product (AGE) accumulation. This study used in vitro, preclinical, and human approaches to explore the potential effects of these agents on the modulation of the receptor for AGE (RAGE). Bovine aortic endothelial cells that were exposed to the angiotensin-converting enzyme inhibitor (ACEi) ramiprilat in the presence of high glucose demonstrated a significant increase in soluble RAGE (sRAGE) secreted into the medium. In streptozotocin-induced diabetic rats, ramipril treatment (ACEi) at 3 mg/L for 24 wk reduced the accumulation of skin collagen-linked carboxymethyllysine and pentosidine, as well as circulating and renal AGE. Renal gene upregulation of total RAGE (all three ~~splice~~ variants) was observed in ACEi-treated animals. There was a specific increase in the gene expression of the ~~splice~~ variant C-truncated RAGE (sRAGE). There were also increases

in sRAGE protein identified within renal cells with ACEi treatment, which showed AGE-binding ability. This was associated with decreases in renal full-length RAGE protein from ACEi-treated rats. Decreases in plasma soluble RAGE that were significantly increased by ACEi treatment were also identified in diabetic rats. Similarly, there was a significant increase in plasma sRAGE in patients who had type 1 diabetes and were treated with the ACEi perindopril. Complexes between sRAGE and carboxymethyllysine were identified in human and rodent diabetic plasma. It is postulated that ACE inhibition reduces the accumulation of AGE in diabetes partly by increasing the production and secretion of sRAGE into plasma.

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18446161 BIOSIS NO.: 200510140661

Detection of secreted exon 8 lacking splice variant of receptor for advanced glycation end products (RAGE) from astrocytes

AUTHOR: Rouhiainen A (Reprint); Kuja-Panula J; Rauvala H

AUTHOR ADDRESS: Univ Helsinki, Ctr Neurosci, Helsinki, Finland**Finland

JOURNAL: Journal of Neurochemistry 90 (Suppl. 1): p61 AUG 04 2004

CONFERENCE/MEETING: 35th Annual Meeting of the

Transactions-of-the-American-Society-for-Neurochemistry New York, NY, USA
August 14 -18, 2004; 20040814

SPONSOR: Transact Amer Soc Neurochem

ISSN: 0022-3042

DOCUMENT TYPE: Meeting; Meeting Poster

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7/7/10

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18336797 BIOSIS NO.: 200510031297

Roles of the receptor for advanced glycation endproducts in diabetes-induced vascular injury

AUTHOR: Yonekura Hideto (Reprint); Yamamoto Yasuhiko; Sakurai Shigeru;
Watanabe Takuo; Yamamoto Hiroshi

AUTHOR ADDRESS: Kanazawa Univ, Grad Sch Med Sci, Dept Biochem and Mol Vasc Biol, Kanazawa, Ishikawa 9208640, Japan**Japan

AUTHOR E-MAIL ADDRESS: hyone@med.kanazawa-u.ac.jp

JOURNAL: Journal of Pharmacological Sciences 97 (3): p305-311 MAR 05 2005

ISSN: 1347-8613

DOCUMENT TYPE: Article; Literature Review

RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: Diabetic patients have shorter life span and poorer Quality of Life mainly due to diabetic vascular complications. Recent in vitro and in vivo studies have shown that advanced glycation endproducts (AGE) account for diabetic vascular complications through their engagement of the receptor for AGE (RAGE). In this review, we summarize our recent studies on the roles of the AGE-RAGE system in diabetes-induced vascular

injury. In vitro experiments showed that AGE engagement of RAGE leads to changes in endothelial cells (EC) and pericytes, which are characteristic of diabetic microangiopathy. Diabetic RAGE transgenic mice that overexpress RAGE in vascular cells exhibited the exacerbation of the indices of nephropathy and retinopathy, and this was prevented by the inhibition of AGE formation. RAGE overexpression also caused calcium handling impairment in cardiac myocytes. In contrast to the RAGE-overexpressing mice, diabetic RAGE knockout mice showed marked improvement of nephropathy. We found that human vascular cells express a novel splice variant coding for a soluble RAGE protein and named it endogenous secretory RAGE (esRAGE). The esRAGE neutralizes AGE actions on EC and is present in human sera. Individual variations in circulating esRAGE could be a determinant for individual differences in susceptibility or resistance to the development of diabetic vascular complications. The AGE-RAGE system should be, therefore, a candidate molecular target for overcoming diabetic vascular complications.

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17915973 BIOSIS NO.: 200400286730

Expression of a novel secreted splice variant of the receptor for advanced glycation end products (RAGE) in human brain astrocytes and peripheral blood mononuclear cells

AUTHOR: Youn Ju Ho (Reprint); Park In Ho; Yeon Soo In; Choi In-Hong; Shin Jeon-Soo

AUTHOR ADDRESS: Department of Microbiology, BK21 project for Medical Science, Yonsei University College of Medicine, 134 Shinchon-dong Seodaemun-gu, Seoul, 120-752, South Korea**South Korea

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JOURNAL: FASEB Journal 18 (4-5): pAbst. 331.28 2004 2004

MEDIUM: e-file

CONFERENCE/MEETING: FASEB Meeting on Experimental Biology: Translating the Genome Washington, District of Columbia, USA April 17-21, 2004; 20040417

SPONSOR: FASEB

ISSN: 0892-6638 (ISSN print)

DOCUMENT TYPE: Meeting; Meeting Abstract

RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: The engagement of the receptor for advanced glycation end products (RAGE) on the cell surface induces cellular dysfunction in a number of pathophysiological situations of vascular dysfunction, tumor cell invasion, inflammatory response, and T cell infiltration. The administration of truncated, soluble RAGE can modulate RAGE-mediated perturbations. Here, we report a novel splice variant (Δ 8-RAGE) of RAGE mRNA, which lacks exon 8 of the genomic RAGE gene and contains an early stop codon in exon 10 due to a frame shift mutation. Δ 8-RAGE mRNA was found in human primary astrocytes and peripheral blood mononuclear cells (PBMCs). Transient transfection experiments demonstrated that Δ 8-RAGE mRNA was translated into a secretory protein as deduced. Moreover, two different segments of the spliced variant were identified in PBMCs by RT-PCR. The findings of this study suggest that the diverse splicing variants of RAGE are possible in many tissues and their products may influence the RAGE-mediated

pathogenesis and immune modulation.

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DIALOG(R)File 5:Biosis Previews(R)

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Expression of a novel secreted ~~splice~~ variant of the ~~receptor~~ for ~~advanced~~ ~~glycation~~ end products (RAGE) in human brain astrocytes and peripheral blood mononuclear cells.

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RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: The engagement of the ~~receptor~~ for ~~advanced~~ ~~glycation~~ end products (RAGE) on the cell surface induces cellular dysfunction in a number of pathophysiological situations of vascular dysfunction, tumor cell invasion, inflammatory response, and T cell infiltration. The administration of truncated, soluble RAGE can modulate RAGE-mediated perturbations. Here, we report a novel ~~splice~~ variant (DELTA8-RAGE) of RAGE mRNA, which lacks exon 8 of the genomic RAGE gene and contains an early stop codon in exon 10 due to a frame shift mutation. DELTA8-RAGE mRNA was found in human primary astrocytes and peripheral blood mononuclear cells (PBMCs). Transient transfection experiments demonstrated that DELTA8-RAGE mRNA was translated into a secretory protein as deduced. Moreover, two different segments of the spliced variant were identified in PBMCs by RT-PCR. The findings of this study suggest that the diverse splicing variants of RAGE are possible in many tissues and their products may influence the RAGE-mediated pathogenesis and immune modulation.

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Tissue-specific expression patterns of the RAGE receptor and its soluble forms: A result of regulated alternative splicing?

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ABSTRACT: The receptor for advanced glycation end products (RAGE) is known to be causally involved in a variety of pathophysiological processes, e.g. immune/inflammatory disorders, Alzheimer disease, tumors, and abnormalities associated with diabetes as arteriosclerosis or disordered wound healing. So far, human cDNAs have been characterized encoding for the RAGE receptor and a truncated soluble form lacking the transmembrane and the cytosolic domain. The latter form represents a naturally occurring competitive inhibitor of signalling pathways induced by the membrane-standing RAGE receptor. In order to perform a relative expression analysis of both RAGE forms, an RT-PCR experiment was designed allowing the simultaneous amplification of corresponding transcripts. We were able to identify three novel human RAGE transcripts all encoding truncated soluble forms of RAGE. The relative expression ratios for the full-length RAGE transcript to the sum of its splice-variants encoding the soluble variants varied strongly among the tissues tested. Therefore, the pre-mRNA of RAGE must be subject to regulated alternative splicing activated by extracellular cues of yet unknown cellular signalling pathways. Thus, as deduced from the occurrence at the RNA level, it can be hypothesized that there is a complex RAGE regulation network involving isoforms competing for the binding of ligands.

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17238754 BIOSIS NO.: 200300197473

Novel splice variants of the receptor for advanced glycation end-products expressed in human vascular endothelial cells and pericytes, and their putative roles in diabetes-induced vascular injury.

AUTHOR: Yonekura Hideto; Yamamoto Yasuhiko; Sakurai Shigeru; Petrova Ralica G; Abedin Md Joynal; Li Hui; Yasui Kiyoshi; Takeuchi Masayoshi; Makita Zenji; Takasawa Shin; Okamoto Hiroshi; Watanabe Takuo; Yamamoto Hiroshi (Reprint)

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LANGUAGE: English

ABSTRACT: The binding of advanced glycation end-products (AGE) to the receptor for AGE (RAGE) is known to deteriorate various cell functions and is implicated in the pathogenesis of diabetic vascular complications. In the present study, we show that the cellular constituents of small vessels, endothelial cells (EC) and pericytes express novel splice

variants of RAGE mRNA coding for the isoforms that lack the N-terminal V-type immunoglobulin-like domain (N-truncated) or the C-terminal transmembrane domain (C-truncated), as well as the known full-length mRNA. The ratio of the expression of the three variants was different between EC and pericytes; the content of the C-truncated form was highest in EC, whereas the full-length form was the most abundant in pericytes. Transfection experiments with COS-7 cells demonstrated that those variant mRNAs were translated into proteins as deduced; C-truncated RAGE was efficiently secreted into the culture media, and N-truncated RAGE was located mainly on the plasma membrane. The three isoforms were also detected in primary cultured human EC and pericytes. Further, full-length and C-truncated forms of RAGE bound to an AGE-conjugated column, whereas N-truncated RAGE did not. The AGE induction of extracellular-signal-related kinase phosphorylation and vascular endothelial growth factor in EC and of the growth and cord-like structure formation of EC was abolished completely by C-truncated RAGE, indicating that this endogenous secretory receptor (endogenous secretory RAGE) is cytoprotective against AGE. The results may contribute to our understanding of the molecular basis for the diversity of cellular responses to AGE and for individual variations in the susceptibility to diabetic vascular complications.

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